SEQUENCE LISTING

GENERAL INFORMATION:

(i) APPLICANT: Roncarolo, Maria-Grazia de Waal Malefyt, Rene Bacchetta, Rosa Groux, Herve M. de Vries, Jan E.

- (ii) TITLE OF INVENTION: USE OF INTERLEUKIN-10 TO PRODUCE A POPULATION OF SUPPRESSOR CELLS
- (iii) NUMBER OF SEQUENCES: 25
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: DNAX Research Institute
 - (B) STREET: 901 California Avenue
 - (C) CITY: Palo Alto
 - (D) STATE: California
 - (E) COUNTRY: USA
 - (F) ZIP: 94304-1104
 - (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: Apple Macintosh
 - (C) OPERATING SYSTEM: Macintosh 7.5.5
 - (D) SOFTWARE: Microsoft Word 5.1a
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 08/643,810
 - (B) FILING DATE: 06-MAY-1996
 - (C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 07/846,208
 - (B) FILING DATE: 04-MAR-1992
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Ching, Edwin P.
 - (B) REGISTRATION NUMBER: 34,090
 - (C) REFERENCE/DOCKET NUMBER: DX0261K1
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: 415-852-9196
 - (B) TELEFAX: 415-496-1200
- (2) INFORMATION FOR SEQ ID NO:1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 178 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

Met His Ser Ser Ala Leu Leu Cys Cys Leu Val Leu Leu Thr Gly Val 1 5 10 15

Arg Ala Ser Pro Gly Gln Gly Thr Gln Ser Glu Asn Ser Cys Thr His 20 25 30

Phe Pro Gly Asn Leu Pro Asn Met Leu Arg Asp Leu Arg Asp Ala Phe 35 40 45

Ser Arg Val Lys Thr Phe Phe Gln Met Lys Asp Gln Leu Asp Asn Leu 50 55 60

Leu Leu Lys Glu Ser Leu Leu Glu Asp Phe Lys Gly Tyr Leu Gly Cys 65 70 75 80

Gln Ala Leu Ser Glu Met Ile Gln Phe Tyr Leu Glu Glu Val Met Pro 85 90 95

Gln Ala Glu Asn Gln Asp Pro Asp Ile Lys Ala His Val Asn Ser Leu 100 105 110

Gly Glu Asn Leu Lys Thr Leu Arg Leu Arg Leu Arg Arg Cys His Arg 115 120 125

Phe Leu Pro Cys Glu Asn Lys Ser Lys Ala Val Glu Gln Val Lys Asn 130 135 140

Ala Phe Asn Lys Leu Gln Glu Lys Gly Ile Tyr Lys Ala Met Ser Glu
145 150 155 160

Phe Asp Ile Phe Ile Asn Tyr Ile Glu Ala Tyr Met Thr Met Lys Ile 165 170 175

Arg Asn

(2) INFORMATION FOR SEQ ID NO:2:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 170 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

Met Glu Arg Arg Leu Val Val Thr Leu Gln Cys Leu Val Leu Tyr

1				5					10			,		15	
Leu	Ala	Pro	Glu 20	Cys	Gly	Gly	Thr	Asp 25	Gln	Cys	Asp	Asn	Phe 30	Pro	Gln
Met	Leu	Arg 35	Asp	Leu	Arg	Asp	Ala 40	Phe	Ser	Arg	Val	Lys 45	Thr	Phe	Phe
Gln	Thr 50	Lys	Asp.	Glu	Val	Asp 55	Asn	Leu	Leu	Leu	Lys 60	Glu	Ser	Leu	Leu
Gļu 65	Asp	Phe	Lys	Gly	Tyr 70	Leu	Gly	Cys	Gln	Ala 75	Leu	Ser	Glu	Met	Ile 80
Gln	Phe	Tyr	Leu	Glu 85	Glu	Val	Met	Pro	Gln 90	Ala	Glu	Asn	Gln	Asp 95	Pro
Glu	Ala	Lys	Asp 100	His	Val	Asn	Ser	Leu 105	Gly	Glu	Asn	Leu	Lys 110	Thr	Leu
Arg	Leu	Arg 115	Leu	Arg	Arg	Cys	His 120	Arg	Phe	Leu	Pro	Cys 125	Glu	Asn	Lys
Ser	Lys 130	Ala	Val	Glu	Gln	Ile 135	Lys	Asn	Ala	Phe	Asn 140	Lys	Leu	Gln	Glu
Lys 145	Gly	Ile	Tyr	Lys	Ala 150	Met	Ser	Glu	Phe	Asp 155	Ile	Phe	Ile	Asn	Tyr 160
Ile	Glu	Ala	Tyr	Met 165		Ile	Lys	Ala	Arg 170						
INFORMATION FOR SEQ ID NO:3:															

(2)

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 160 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

Ser Pro Gly Gln Gly Thr Gln Ser Glu Asn Ser Cys Thr His Phe Pro

Gly Asn Leu Pro Asn Met Leu Arg Asp Leu Arg Asp Ala Phe Ser Arg

Val Lys Thr Phe Phe Gln Met Lys Asp Gln Leu Asp Asn Leu Leu

Lys Glu Ser Leu Leu Glu Asp Phe Lys Gly Tyr Leu Gly Cys Gln Ala 55

Leu Ser Glu Met Ile Gln Phe Tyr Leu Glu Glu Val Met Pro Gln Ala 65 70 75 80

Glu Asn Gln Asp Pro Asp Ile Lys Ala His Val Asn Ser Leu Gly Glu 85 90 95

Asn Leu Lys Thr Leu Arg Leu Arg Leu Arg Arg Cys His Arg Phe Leu 100 105 110

Pro Cys Glu Asn Lys Ser Lys Ala Val Glu Gln Val Lys Asn Ala Phe 115 120 125

Asn Lys Leu Gln Glu Lys Gly Ile Tyr Lys Ala Met Ser Glu Phe Asp 130 135 140

Ile Phe Ile Asn Tyr Ile Glu Ala Tyr Met Thr Met Lys Ile Arg Asn 145 150 155 160

(2) INFORMATION FOR SEQ ID NO:4:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 147 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

Thr Asp Gln Cys Asp Asn Phe Pro Gln Met Leu Arg Asp Leu Arg Asp 1 10 15

Ala Phe Ser Arg Val Lys Thr Phe Phe Gln Thr Lys Asp Glu Val Asp 20 25 30

Asn Leu Leu Lys Glu Ser Leu Leu Glu Asp Phe Lys Gly Tyr Leu 35 40 45

Gly Cys Gln Ala Leu Ser Glu Met Ile Gln Phe Tyr Leu Glu Glu Val 50 55 60

Met Pro Gln Ala Glu Asn Gln Asp Pro Glu Ala Lys Asp His Val Asn 65 70 75 80

Ser Leu Gly Glu Asn Leu Lys Thr Leu Arg Leu Arg Leu Arg Cys 85 90 95

His Arg Phe Leu Pro Cys Glu Asn Lys Ser Lys Ala Val Glu Gln Ile 100 105 110

Lys Asn Ala Phe Asn Lys Leu Gln Glu Lys Gly Ile Tyr Lys Ala Met 115 120 125

	Ser	Glu P 130	he Asp	Ile Phe	Ile 135	Asn	Tyr ·	Ile	Glu	Ala 140	Tyr	Met	Thr	Ile	
	Lys 145	Ala A	ırg												
(2)	INFO	RMATIC	N FOR	SEQ ID N	0:5:										
	(i)	(A) (B) (C)	LENGTH TYPE: STRAND	ARACTERI: 15 bas nucleic EDNESS: GY: line	e pai acid singl	rs									
	(ii)	MOLEC	CULE TY	PE: cDNA											
	(xi)	SEQUE	ENCE DE	CSCRIPTIO	N: SE	EQ I	D NO	:5:							
CTAA	AGGAG(GT TTA	AAC												15
(2)	INFO	RMATIC	ON FOR	SEQ ID N	10:6:										
	(i)	(A) (B) (C)	LENGTH TYPE: STRANE	HARACTERI H: 10 bas nucleic DEDNESS: DGY: line	e pai acid singl	irs									
	(ii)	MOLE	CULE TY	PE: cDNA											
	(xi)	SEQU	ENCE DE	ESCRIPTIO	N: SI	EQ I	D NO	:6:							
ATG	AGCTC	AΤ													10
(2)	INFO	RMATI(ON FOR	SEQ ID N	10:7:										
	(i)	(A) (B) (C)	LENGTI TYPE: STRANI	HARACTER: H: 60 bas nucleic DEDNESS: DGY: line	se pa acid sing	irs									
	(ii)	MOLE	CULE T	YPE: cDN	A										
	(xi)	SEQU	ENCE D	ESCRIPTION TO SERVICE	ON: S	EQ I	D NC	:7:							

AGCCCAGGCC AGGGCACCCA GTCTGAGAAC AGCTGCACCC ACTTCCCAGG TAACCGGTAC 60

(2)	INFORMATION FOR SEQ ID NO:8:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 56 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(ii) MOLECULE TYPE: cDNA	
	• · · · · · · · · · · · · · · · · · · ·	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:	
CGGT	TTACCTG GGAAGTGGGT GCAGCTGTTC TCAGACTGGG TGCCCTGGCC TGGGCT	56
(2)	INFORMATION FOR SEQ ID NO:9:	
	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 62 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: cDNA	
СТА	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9: ACCTGCC TAACATGCTT CGAGATCTCC GAGATGCCTT CAGCAGAGTG AAGACTTTCT	60
тт	Accross Transfer Comments	62
	INFORMATION FOR SEQ ID NO:10:	
(2)	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 48 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: cDNA	
		•
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:	
CTT	CACTCTG CTGAAGGCAT CTCGGAGATC TCGAAGCATG TTAGGCAG	48
(2)	INFORMATION FOR SEQ ID NO:11:	
	(i) SEQUENCE CHARACTERISTICS:	

(A) LENGTH: 35 base pairs

	(ii) MOLECULE TYPE: cDNA	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:	
		35
CAA	ATGĄAGG ATCAGCTGGA CAACTTGTTC TTAAG	33
(2)	INFORMATION FOR SEQ ID NO:12:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 44 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(ii) MOLECULE TYPE: cDNA	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:	
CTT	AAGAACA AGTTGTCCAG CTGATCCTTC ATTTGAAAGA AAGT	44
(2)	INFORMATION FOR SEQ ID NO:13:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 69 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(ii) MOLECULE TYPE: cDNA	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:	
GAG	STECTIGE TGGAGGACTT TAAGGGTTAC CTGGGTTGCC AAGCCTTGTC TGAGATGATC	60
CAG	STTTTAT	69
(2)	INFORMATION FOR SEQ ID NO:14:	
	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 73 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: cDNA	

(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:	
GATCATCTCA GACAAGGCTT GGCAACCCAG GTAACCCTTA AAGTCCTCCA GCAAGGACTC	60
CTAGATAAAA CTG	73
(2) INFORMATION FOR SEQ ID NO:15:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 61 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:	
CTAGAGGAGG TGATGCCCCA AGCTGAGAAC CAAGACCCAG ACATCAAGGC GCATGTTAAC	60
G	61
(2) INFORMATION FOR SEQ ID NO:16:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 65 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:	
GTTAACATGC GCCTTGATGT CTGGGTCTTG GTTCTCAGCT TGGGGCATCA CCTCCTCTAG	60
TCGAC	65
(2) INFORMATION FOR SEQ ID NO:17:	
(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 63 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: cDNA	

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:	
AACTCCCTGG GGGAGAACCT GAAGACCCTC AGGCTGAGGC TACGGCGCTG TCATCGATCT	60
GCA	63
(2) INFORMATION FOR SEQ ID NO:18:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 59 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:	
GATCGATGAC AGCGCCGTAG CCTCAGCCTG AGGGTCTTCA GGTTCTCCCC CAGGGAGTT	59
(2) INFORMATION FOR SEQ ID NO:19:	
(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 60 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:	
CGATTTCTTC CCTGTCAAAA CAAGAGCAAG GCCGTGGAGC AGGTGAAGAA CGCGTGCATG	6
CGAT-TTCTTC CCTGTCAAAA CAAGAGCAAG GCCGTGGAGC AGGTGALC.11 GGGGTGTTT	
(2) INFORMATION FOR SEQ ID NO:20:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 54 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(ii) MOLECULE TYPE: cDNA	

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:

CACGCGTTCT TCACCTGCTC CACGGCCTTG CTCTTGTTTT GACAGGGAAG AAAT	54
(2) INFORMATION FOR SEQ ID NO:21:	
(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 58 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:	
CGCGTTTAAT AATAAGCTCC AAGACAAAGG CATCTACAAA GCCATGAGTG AGTTTGAC	58
(2) INFORMATION FOR SEQ ID NO:22:	
(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 45 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:	
ACTCATGGCT TTGTAGATGC CTTTGTCTTG GAGCTTATTA TTAAA	45
(2) INFORMATION FOR SEQ ID NO:23:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 51 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:23:	
ATCTTCATCA ACTACATAGA AGCCTACATG ACAATGAAGA TACGAAACTG A	51
(2) INFORMATION FOR SEQ ID NO:24:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 64 base pairs	

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:24:	
AGCTTCAGTT TCGTATCTTC ATTGTCATGT AGGCTTCTAT GTAGTTGATG AAGATGTCAA	60
ACTC	64
(2) INFORMATION FOR SEQ ID NO:25:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 498 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:25:	
AATTCATGGA GCGAAGGTTA GTGGTCACTC TGCAGTGCCT GGTGCTGCTT TACCTGGCAC	60
CTGAGTGTGG AGGTACAGAC CAATGTGACA ATTTTCCCCA GACCTAAGAG ATGCCTTCAG	120
TCGTGTTAAA ACCTTTTCC AGACAAAGGA CGAGGTAGAT AACCTTTTGC TCAAGGAGTC	180
TCTGCTAGAG GACTTTAAGG ATGCCAGGCC CTGTCAGAAA TGATCCAATT CTACCTGGAG	240
GAAGTCATGC CACAGGCTGA AACCAGGACC CTGAAGCCAA AGACCATGTC AATTCTTTGG	300
GTGAAAATCT AAAGACCCTA CGGCTCCGCC TGCGCAGGTG CCACAGGTTC CTGCCGTGTG	360
AGAACAAGAG TAAAGCTGTG GAACAGATAA AAAATGCCTT TAACAAGCTG CAGGAAAAAG	420
GAATTTACAA AGCCATGAGT GAATTTGACA TTTTTATTAA CTACATAGAA GCATACATGA	480
CAATTAAAGC CAGGTGAG	498
(2) INFORMATION FOR SEQ ID NO:26:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 40 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(ii) MOLECULE TYPE: cDNA	

(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:26:
GAATTCCCAG GTTTAACGTA AGGAGGTTTA ACCATCGATG

40